RAW SEQUENCE LISTING   DATE: 07/17/2001
PATENT APPLICATION: US/09/815,330   TIME: 09:09:05

Input Set : **A:\PTO.txt**
Output Set: **N:\CRF3\07172001\I815330.raw**

Does Not Comply
Corrected Diskette Needed

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      1 <110> APPLICANT: Zaia, John Hawkins,
W-->  2 <120> TITLE OF INVENTION: Protein Kinase Deficient, Immunologically Active CMVpp65
W-->  3 <130> FILE REFERENCE: 1954-333<140> US 09/815,330<141> 2001-03-23<150> US
C-->  4 <140> CURRENT APPLICATION NUMBER: US/09/815,330
C-->  4 <141> CURRENT FILING DATE: 2001-03-23
W-->  4 <151> PRIOR FILING DATE: 2000-03-23<160> 8 <170> PatentIn version 3.0<210> 1<211>
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ERRORED SEQUENCES

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E--> 10 1                    5
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See next page

insert a hard return after each response

SEQUENCE LISTING<110> Zaia, John Hawkins,
Ghislaine<120> Protein Kinase Deficient, Immunologically Active CMVpp65
Mutants<130> 1954-333<140> US 09/815,330<141> 2001-03-23<150> US
60/191,464<151> 2000-03-23<160> 8 <170> PatentIn version 3.0<210> 1<211>
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X = any amino acid<220><221> VARIANT<222> (4)..(4)<223> X = any amino
acid<220><221> VARIANT<222> (5)..(5)<223> X = any amino acid<400> 1

Glu Xaa Glu Xaa Xaa Glu
1   5

major format error - please contact

Robert Wax

at 703-306-4119

or 703-308-4216

for assistance

Also, see cover sheet

FYI Review the Sequence Listing to insure a corresponding
explanation is presented in the <220> to <223> fields of
each sequence using n or Xaa.

Input Set : **A:\PTO.txt**
Output Set: **N:\CRF3\07172001\I815330.raw**

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L:2  M:283 W: Missing Blank Line separator, <120> field identifier
L:3  M:283 W: Missing Blank Line separator, <130> field identifier
L:4  M:270 C: Current Application Number differs, Replaced Current Application No
L:4  M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:4  M:289 W: Identifier Missing or Out-Of-Order, <150> PRIOR APP NO
L:6  M:283 W: Missing Blank Line separator, <220> field identifier
L:6  M:256 W: Invalid Numeric Header Field, <220> has non-blank data
L:7  M:283 W: Missing Blank Line separator, <220> field identifier
L:7  M:256 W: Invalid Numeric Header Field, <220> has non-blank data
L:9  M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:0
L:9  M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:0
L:9  M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:0
L:9  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:0
L:10 M:252 E: No. of Seq. differs, <211>LENGTH:Input:0 Found:6 SEQ:0
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